

# NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

- ☐ 1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
- ☐ 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
- ☐ 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
- ☒ 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
- ☐ 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
- ☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
- ☐ 7. Other: \_\_\_\_\_

## **Applicant Must Provide:**

- ☒ An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
- ☒ An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
- ☒ A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).

For questions regarding compliance to these requirements, please contact:

For Rules Interpretation, call (703) 308-4216

For CRF Submission Help, call (703) 308-4212

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PAGE: 1

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/490,208

DATE: 02/15/2000  
TIME: 15:06:25

Input Set: I490208.RAW

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

Does Not Comply  
Corrected Diskette Needed

1 <110> C. Frank Bennett  
2 Nicholas M. Dean  
3 Lex M. Cowser  
4 <120> ANTISENSE MODULATION OF INDUCIBLE NITRIC OXIDE SYNTHASE EXPRESSION  
5 <130> RTS-0066  
6 <140> US/09/490,208  
7 <141> 2000-01-24  
8 <160> 182

ERRORED SEQUENCES FOLLOW

E--> 9 <210> 10  
10 <211> 4110 4/45 done (p6)  
11 <212> DNA  
12 <213> Mus musculus  
13 <220>  
14 <221> mRNA  
15 <222> (1)...(4110)  
16 <400> 10  
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18 aagttctcaa ggcacaggte tcttctcggt ttgactgtcc ttaccccgagg gaggcagtg 120  
19 agccagctgc aagccccaca gtgaagaaca tctgagctca aatccagata agtgacataa 180  
20 gtgacctgct ttgtaaagcc atagag atg gcc tgt cct tgg aaa ttt ctg ttc 233  
21 Met Ala Cys Pro Trp Lys Phe Leu Phe  
22 1 5  
23 aag acc aaa ttc cac cag tat gca atg aat ggg gaa aaa gac atc aac 281  
24 Lys Thr Lys Phe His Gln Tyr Ala Met Asn Gly Glu Lys Asp Ile Asn  
25 10 15 20 25  
26 aac aat gtg gag aaa gcc ccc tgt gcc acc tcc agt cca gtg aca cag 329  
27 Asn Asn Val Glu Lys Ala Pro Cys Ala Thr Ser Ser Pro Val Thr Gln  
28 30 35 40  
29 gat gac ott cag tat cac aac ctc agc aag cag cag aat gag tcc ccg 377  
30 Asp Asp Leu Gln Tyr His Asn Leu Ser Lys Gln Gln Asn Glu Ser Pro  
31 45 50 55  
32 cag ccc ctc gtg gag acg gga aag aag tct cca gaa tct ctg gtc aag 425  
33 Gln Pro Leu Val Glu Thr Gly Lys Lys Ser Pro Glu Ser Leu Val Lys  
34 60 65 70  
35 ctg gat gca acc cca ttg tcc tcc cca cgg cat gtg agg atc aaa aac 473  
36 Leu Asp Ala Thr Pro Leu Ser Ser Pro Arg His Val Arg Ile Lys Asn

PAGE: 2

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/490,208

DATE: 02/15/2000  
TIME: 15:06:25

Input Set: I490208.RAW

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44	ccc aaa agt ttg acc aga gga ccc agg gac aag cct acc cct cca gat				617
45	Pro Lys Ser Leu Thr Arg Gly Pro Arg Asp Lys Pro Thr Pro Pro Asp				
46		125	130	135	
47	gag ctt cta cct caa gct atc gaa ttt gtc aac caa tat tac ggc tcc				665
48	Glu Leu Leu Pro Gln Ala Ile Glu Phe Val Asn Gln Tyr Tyr Gly Ser				
49		140	145	150	
50	ttc aaa gag gca aaa ata gag gaa cat ctg gcc agg gtg gaa gcg gta				713
51	Phe Lys Glu Ala Lys Ile Glu Glu His Leu Ala Arg Val Glu Ala Val				
52		155	160	165	
53	aca aag gag ata gaa aca aca gga acc tac caa ctg acg gga gat gag				761
54	Thr Lys Glu Ile Glu Thr Thr Gly Thr Tyr Gln Leu Thr Gly Asp Glu				
55		170	175	180	185
56	ctc atc ttc gcc acc aag cag gcc tgg cgc aat gcc cca cgc tgc att				809
57	Leu Ile Phe Ala Thr Lys Gln Ala Trp Arg Asn Ala Pro Arg Cys Ile				
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59	ggg agg atc cag tgg tcc aac ctg cag gtc ttc gat gcc cgc agc tgt				857
60	Gly Arg Ile Gln Trp Ser Asn Leu Gln Val Phe Asp Ala Arg Ser Cys				
61		205	210	215	
62	tcc act gcc cgg gaa atg ttt gaa cac atc tgc aga cac gtg cgt tac				905
63	Ser Thr Ala Arg Glu Met Phe Glu His Ile Cys Arg His Val Arg Tyr				
64		220	225	230	
65	tcc acc aac aat ggc aac atc agg tgc gcc atc acc gtg ttc ccc cag				953
66	Ser Thr Asn Asn Gly Asn Ile Arg Ser Ala Ile Thr Val Phe Pro Gln				
67		235	240	245	
68	cgg agt gat ggc aag cac gac ttc cgg gtg tgg aat gct cag ctc atc				1001
69	Arg Ser Asp Gly Lys His Asp Phe Arg Val Trp Asn Ala Gln Leu Ile				
70		250	255	260	265
71	cgc tat gct ggc tac cag atg cca gat ggc agc atc aga ggg gac cct				1049
72	Arg Tyr Ala Gly Tyr Gln Met Pro Asp Gly Ser Ile Arg Gly Asp Pro				
73		270	275	280	
74	gcc aac gtg gaa ttc act cag ctg tgc atc gac ctg ggc tgg aag ccc				1097
75	Ala Asn Val Glu Phe Thr Gln Leu Cys Ile Asp Leu Gly Trp Lys Pro				
76		285	290	295	
77	aag tac ggc cgc ttc gat gtg gtc ccc ctg gtc ctg cag gcc aat ggc				1145
78	Lys Tyr Gly Arg Phe Asp Val Val Pro Leu Val Leu Gln Ala Asn Gly				
79		300	305	310	
80	cgt gac cct gag ctc ttc gaa atc cca cct gac ctt gtg ctt gag gtg				1193
81	Arg Asp Pro Glu Leu Phe Glu Ile Pro Pro Asp Leu Val Leu Glu Val				
82		315	320	325	
83	gcc atg gaa cat ccc aaa tac gag tgg ttt cgg gaa ctg gag cta aag				1241
84	Ala Met Glu His Pro Lys Tyr Glu Trp Phe Arg Glu Leu Glu Leu Lys				
85		330	335	340	345
86	tgg tac gcc ctg cct gca gtg gcc aac atg ctg ctt gag gtg ggc ggc				1289
87	Trp Thr Ala Thr Pro Phe Thr Thr Thr Thr Thr Thr Thr Thr Thr Thr				

PAGE: 3

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION US/09/490,208**

 DATE: 02/15/2000  
 TIME: 15:06:25

Input Set: I490208.RAW

90	Leu Glu Phe Pro Gly Cys Pro Phe Asn Gly Trp Tyr Met Gly Thr Glu	
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93	Ile Gly Val Arg Asp Phe Cys Asp Val Gln Arg Tyr Asn Ile Leu Glu	
94	380 385 390	
95	gaa gtg ggc agg aga atg ggc ctg gaa acg cac aag ctg gcc tcg ctc	1433
96	Glu Val Gly Arg Arg Met Gly Leu Glu Thr His Lys Leu Ala Ser Leu	
97	395 400 405	
98	tgg aaa gac cag gct gtc gtt gag atc aac att gct gtg atc cat agt	1481
99	Trp Lys Asp Gln Ala Val Val Glu Ile Asn Ile Ala Val Ile His Ser	
100	410 415 420 425	
101	ttt cag aag cag aat gtg acc atc atg gac cac cac tcg gct gca gaa	1529
102	Phe Gln Lys Gln Asn Val Thr Ile Met Asp His His Ser Ala Ala Glu	
103	430 435 440	
104	tcc ttc atg aag tac atg cag aat gaa tac cgg tcc cgt ggg ggc tgc	1577
105	Ser Phe Met Lys Tyr Met Gln Asn Glu Tyr Arg Ser Arg Gly Gly Cys	
106	445 450 455	
107	ccg gca gac tgg att tgg ctg gtc cct ccc atg tct ggg agc atc acc	1625
108	Pro Ala Asp Trp Ile Trp Leu Val Pro Pro Met Ser Gly Ser Ile Thr	
109	460 465 470	
110	ccc gtg ttt cac cag gag atg ctg aac tac gtc ctg tcc cct ttc tac	1673
111	Pro Val Phe His Gln Glu Met Leu Asn Tyr Val Leu Ser Pro Phe Tyr	
112	475 480 485	
113	tac tat cag gta gag gcc tgg aaa acc cat gtc tgg cag gac gag aag	1721
114	Tyr Tyr Gln Val Glu Ala Trp Lys Thr His Val Trp Gln Asp Glu Lys	
115	490 495 500 505	
116	cgg aga ccc aag aga aga gag att cca ttg aaa gtc ttg gtc aaa gct	1769
117	Arg Arg Pro Lys Arg Arg Glu Ile Pro Leu Lys Val Leu Val Lys Ala	
118	510 515 520	
119	gtg ctc ttt gcc tgt atg ctg atg cgc aag aca atg gcg tcc cga gtc	1817
120	Val Leu Phe Ala Cys Met Leu Met Arg Lys Thr Met Ala Ser Arg Val	
121	525 530 535	
122	aga gtc acc atc ctc ttt gcg aca gag aca gga aaa tca gag gcg ctg	1865
123	Arg Val Thr Ile Leu Phe Ala Thr Glu Thr Gly Lys Ser Glu Ala Leu	
124	540 545 550	
125	gcc tgg gac ctg ggg gcc tta ttc agc tgt gcc ttc aac ccc aag gtt	1913
126	Ala Trp Asp Leu Gly Ala Leu Phe Ser Cys Ala Phe Asn Pro Lys Val	
127	555 560 565	
128	gtc tgc atg gat aag tac agg ctg agc tgc ctg gag gag gaa cgg ctg	1961
129	Val Cys Met Asp Lys Tyr Arg Leu Ser Cys Leu Glu Glu Glu Arg Leu	
130	570 575 580 585	
131	ctg ttg gtg gtg acc agt acg ttt ggc aat gga gac tgc cct ggc aat	2009
132	Leu Leu Val Val Thr Ser Thr Phe Gly Asn Gly Asp Cys Pro Gly Asn	
133	590 595 600	
134	gga gag aaa ctg aag aaa tcg ctc ttc atg ctg aaa gag ctc aac aac	2057
135	Gly Glu Lys Leu Lys Lys Ser Leu Phe Met Leu Lys Glu Leu Asn Asn	
136	605 610 615	

PAGE: 4

# RAW SEQUENCE LISTING PATENT APPLICATION US/09/490,208

DATE: 02/15/2000  
TIME: 15:06:25

Input Set: I490208.RAW

140	ttc tgc gcc ttt gct cat gac att gat cag aag ctg tcc cac ctg ggg	2153
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142	635 640 645	
143	gcc tct cag ctc acc ccg atg gga gaa ggg gat gag ctc agt ggg cag	2201
144	Ala Ser Gln Leu Thr Pro Met Gly Glu Gly Asp Glu Leu Ser Gly Gln	
145	650 655 660 665	
146	gag gac gcc ttc cgc agc tgg gcc gtg caa acc ttc aag gca gcc tgt	2249
147	Glu Asp Ala Phe Arg Ser Trp Ala Val Gln Thr Phe Lys Ala Ala Cys	
148	670 675 680	
149	gag acg ttt gat gtc cga ggc aaa cag cac att cag atc ccc aag ctc	2297
150	Glu Thr Phe Asp Val Arg Gly Lys Gln His Ile Gln Ile Pro Lys Leu	
151	685 690 695	
152	tac acc tcc aat gtg acc tgg gac ccg cac cac tac agg ctc gtg cag	2345
153	Tyr Thr Ser Asn Val Thr Trp Asp Pro His His Tyr Arg Leu Val Gln	
154	700 705 710	
155	gac tca cag cct ttg gac ctc agc aaa gcc ctc agc agc atg cat gcc	2393
156	Asp Ser Gln Pro Leu Asp Leu Ser Lys Ala Leu Ser Ser Met His Ala	
157	715 720 725	
158	aag aac gtg ttc acc atg agg ctc aaa tct cgg cag aat cta caa agt	2441
159	Lys Asn Val Phe Thr Met Arg Leu Lys Ser Arg Gln Asn Leu Gln Ser	
160	730 735 740 745	
161	ccg aca tcc agc cgt gcc acc atc ctg gtg gaa ctc tcc tgt gag gat	2489
162	Pro Thr Ser Ser Arg Ala Thr Ile Leu Val Glu Leu Ser Cys Glu Asp	
163	750 755 760	
164	ggc caa ggc ctg aac tac ctg ccg ggg gag cac ctt ggg gtt tgc cca	2537
165	Gly Gln Gly Leu Asn Tyr Leu Pro Gly Glu His Leu Gly Val Cys Pro	
166	765 770 775	
167	ggc aac cag ccg gcc ctg gtc caa ggc atc ctg gag cga gtg gtg gat	2585
168	Gly Asn Gln Pro Ala Leu Val Gln Gly Ile Leu Glu Arg Val Val Asp	
169	780 785 790	
170	ggc ccc aca ccc cac cag aca gtg cgc ctg gag gac ctg gat gag agt	2633
171	Gly Pro Thr Pro His Gln Thr Val Arg Leu Glu Asp Leu Asp Glu Ser	
172	795 800 805	
173	ggc agc tac tgg gtc agt gac aag agg ctg ccc ccc tgc tca ctc agc	2681
174	Gly Ser Tyr Trp Val Ser Asp Lys Arg Leu Pro Pro Cys Ser Leu Ser	
175	810 815 820 825	
176	cag gcc ctc acc tac tcc ccg gac atc acc aca ccc cca acc cag ctg	2729
177	Gln Ala Leu Thr Tyr Ser Pro Asp Ile Thr Thr Pro Pro Thr Gln Leu	
178	830 835 840	
179	ctg ctc caa aag ctg gcc cag gtg gcc aca gaa gag cct gag aga cag	2777
180	Leu Leu Gln Lys Leu Ala Gln Val Ala Thr Glu Glu Pro Glu Arg Gln	
181	845 850 855	
182	agg ctg gag gcc ctg tgc cag ccc tca gag tac agc aag tgg aag ttc	2825
183	Arg Leu Glu Ala Leu Cys Gln Pro Ser Glu Tyr Ser Lys Trp Lys Phe	
184	860 865 870	
185	acc aac agc ccc aca ttc ctg gag gtg cta gag gag ttc ccg tcc ctg	2873
186	Thr Asn Ser Pro Thr Phe Leu Glu Val Leu Glu Glu Phe Pro Ser Leu	
187	875 880 885	

PAGE: 5

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/490,208

DATE: 02/15/2000  
TIME: 15:06:25

Input Set: I490208.RAW

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192	Arg Phe Tyr Ser Ile Ser Ser Ser Arg Asp His Thr Pro Thr Glu Ile				
193		910	915	920	
194	cac ctg act gtg gcc gtg gtc acc tac cac acc gga gat ggc cag ggt				3017
195	His Leu Thr Val Ala Val Val Thr Tyr His Thr Gly Asp Gly Gln Gly				
196		925	930	935	
197	ccc ctg cac cac ggt gtc tgc agc aca tgg ctc aac agc ctg aag ccc				3065
198	Pro Leu His His Gly Val Cys Ser Thr Trp Leu Asn Ser Leu Lys Pro				
199		940	945	950	
200	caa gac cca gtg ccc tgc ttt gtg cgg aat gcc agc gcc ttc cac ctc				3113
201	Gln Asp Pro Val Pro Cys Phe Val Arg Asn Ala Ser Ala Phe His Leu				
202		955	960	965	
203	ccc gag gat ccc tcc cat cct tgc atc ctc atc ggg cct ggc aca ggc				3161
204	Pro Glu Asp Pro Ser His Pro Cys Ile Leu Ile Gly Pro Gly Thr Gly				
205		970	975	980	985
206	atc gtg ccc ttc cgc agt ttc tgg cag caa cgg ctc cat gac tcc cag				3209
207	Ile Val Pro Phe Arg Ser Phe Trp Gln Gln Arg Leu His Asp Ser Gln				
208		990	995	1000	
209	cac aag gga gtg cgg gga ggc cgc atg acc ttg gtg ttt ggg tgc cgc				3257
210	His Lys Gly Val Arg Gly Gly Arg Met Thr Leu Val Phe Gly Cys Arg				
211		1005	1010	1015	
212	cgc cca gat gag gac cac atc tac cag gag gag atg ctg gag atg gcc				3305
213	Arg Pro Asp Glu Asp His Ile Tyr Gln Glu Glu Met Leu Glu Met Ala				
214		1020	1025	1030	
215	cag aag ggg gtg ctg cat gcg gtg cac aca gcc tat tcc cgc ctg cct				3353
216	Gln Lys Gly Val Leu His Ala Val His Thr Ala Tyr Ser Arg Leu Pro				
217		1035	1040	1045	
218	ggc aag ccc aag gtc tat gtt cag gac atc ctg cgg cag cag ctg gcc				3401
219	Gly Lys Pro Lys Val Tyr Val Gln Asp Ile Leu Arg Gln Gln Leu Ala				
220		1050	1055	1060	1065
221	agc gag gtg ctc cgt gtg ctc cac aag gag cca ggc cac ctc tat gtt				3449
222	Ser Glu Val Leu Arg Val Leu His Lys Glu Pro Gly His Leu Tyr Val				
223		1070	1075	1080	
224	tgc ggg gat gtg cgc atg gcc cgg gac gtg gcc cac acc ctg aag cag				3497
225	Cys Gly Asp Val Arg Met Ala Arg Asp Val Ala His Thr Leu Lys Gln				
226		1085	1090	1095	
227	ctg gtg gct gcc aag ctg aaa ttg aat gag gag cag gtc gag gac tat				3545
228	Leu Val Ala Ala Lys Leu Lys Leu Asn Glu Glu Gln Val Glu Asp Tyr				
229		1100	1105	1110	
230	ttc ttt cag ctc aag agc cag aag cgc tat cac gaa gat atc ttc ggt				3593
231	Phe Phe Gln Leu Lys Ser Gln Lys Arg Tyr His Glu Asp Ile Phe Gly				
232		1115	1120	1125	
233	gct gta ttt cct tac gag gcg aag aag gac agg gtg gcg gtg cag ccc				3641
234	Ala Val Phe Pro Tyr Glu Ala Lys Lys Asp Arg Val Ala Val Gln Pro				
235		1130	1135	1140	1145
236	agc agc ctg gag atg tca gcg ctc tga gggcctacag gaggggttaa				3688

PAGE: 6

**RAW SEQUENCE LISTING**  
PATENT APPLICATION US/09/490,208DATE: 02/15/2000  
TIME: 15:06:25

Input Set: I490208.RAW

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242	tctctcaaac	tggggcctcc	ctggtcctt	ggagacaaaa	tcttaaatgc	caggcctggc	3928
243	gagtgggtga	aagatggaac	ttgctgctga	gtgcaccact	tcaagtgacc	accaggaggt	3988
244	gctatcgac	cactgtgtat	ttaaactgct	tgtgtacagt	tatttatgcc	tctgtattta	4048
245	aaaaactaac	accagtcctg	ttcccatg	ccacttgggt	cttccctgta	tgattccttg	4108
246	atggagatat	ttacatgaat	tgcattttac	tttaac			4145

↓

Please review the  
Sequence Listing to ensure that a corresponding explanation is presented in the <220> to  
<223> fields of each sequence which presents at least one n or Xaa.

PAGE: 7

VERIFICATION SUMMARY  
PATENT APPLICATION US/09/490,208

DATE: 02/15/2000  
TIME: 15:06:25

Input Set: I490208.RAW

Line ? Error/Warning

Original Text

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10 E Input 4110, Calc# Bases 4145 differ

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<211> 4110